

# SEQUENCE LISTING

<110> OLSON, ERIC  
SPENCER, JEFFREY A.

<120> METHODS AND COMPOSITIONS FOR STABILIZING MICROTUBULES  
IN STRIATED MUSCLE CELLS

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<150> 60/219,020

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<170> PatentIn Ver. 2.1

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Ile Cys Leu Glu Met Phe Thr Lys Pro Val Val Ile Leu Pro Cys Gln	30	35	40	
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Tyr Lys Gln Glu Ser Thr Arg Pro Glu Lys Lys Leu Asp Gln Pro Met	110	115	120	
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Cys Glu Glu His Glu Glu Glu Arg Ile Asn Ile Tyr Cys Leu Asn Cys	125	130	135	
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Leu Ser Gln Lys Phe Asp Thr Leu Tyr Ala Ile Leu Asp Glu Lys Lys
210 215 220

agc gag ctg ctg cag cgg atc acg cag gag cag gag gag aag ctg ggc 1018
Ser Glu Leu Leu Gln Arg Ile Thr Gln Glu Gln Glu Lys Leu Gly
225 230 235 240

ttc atc gag gct ctg atc ctc cag tac agg gag cag ctg gaa aag tcc 1066
Phe Ile Glu Ala Leu Ile Leu Gln Tyr Arg Glu Gln Leu Glu Lys Ser
245 250 255

acc aag ctt gtg gag acc gcc atc cag tcc ctg gat gag ccc gga ggg 1114
Thr Lys Leu Val Glu Thr Ala Ile Gln Ser Leu Asp Glu Pro Gly Gly
260 265 270

gct acc ttc ctc tca agt gcc aag cag ctc atc aag agc att gta gaa 1162
Ala Thr Phe Leu Ser Ser Ala Lys Gln Leu Ile Lys Ser Ile Val Glu
275 280 285

gcc tcc aag ggc tgc cag ctg ggg aag aca gag caa ggc ttt gag aac 1210
Ala Ser Lys Gly Cys Gln Leu Gly Lys Thr Glu Gln Gly Phe Glu Asn
290 295 300

atg gac tac ttt act ctg gac tta gaa cac ata gca gag gcc ttg agg 1258
Met Asp Tyr Phe Thr Leu Asp Leu Glu His Ile Ala Glu Ala Leu Arg
305 310 315 320

gcc att gac ttt ggg aca ggt aaa gga tgt gat gtt aca tgt ttg acc 1306
Ala Ile Asp Phe Gly Thr Gly Lys Gly Cys Asp Val Thr Cys Leu Thr
325 330 335

ttt gaa agg cag cgt tcc tct tgagttctga ggggaactgt taaaaaagtc 1357
Phe Glu Arg Gln Arg Ser Ser
340

aaatttacac agccagtgtt gacaggtctc tctatggagc cctgactgtc ttagtagtgt 1417

ctaagtagac caagctggtc tggaacacat agagatctat cttgcccatc tctgcttctt 1477

gagggatgag ataaaaggca tgtgcccacc atgcctggct ccacagacaa ctttgtgatg 1537

gatccagggt ctggcacagt gcctggtaca taattgtttc gaaataaatt atctcgtgcc 1597

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[illegible]

Met Asp Tyr Lys Ser Ser Leu Ile Pro Asp Gly Asn Ala Met Glu Asn  
1 5 10 15

Pro Val Val Ile Leu Pro Cys Gln His Asn Leu Cys Arg Lys Cys Ala  
35 40 45

Ser Val Ser Met Ser Gly Gly Arg Phe Arg Cys Pro Ser Cys Arg His  
65 70 75 80

Leu Val Glu Asn Ile Ile Asp Ile Tyr Lys Gln Glu Cys Ser Ser Arg  
100 105 110

Pro Leu Gln Lys Gly Ser His Pro Met Cys Lys Glu His Glu Asp Glu  
115 120 125

Lys Ile Asn Ile Tyr Cys Leu Thr Cys Glu Val Pro Thr Cys Ser Leu  
130 135 140

Cys Lys Val Phe Gly Ala His Gln Ala Cys Glu Val Ala Pro Leu Gln  
145 150 155 160

Ser Ile Phe Gln Gly Gln Lys Thr Glu Leu Ser Asn Cys Ile Ser Met  
165 170 175

Leu Val Ala Gly Asn Asp Arg Val Gln Thr Ile Ile Ser Gln Leu Glu  
180 185 190

Asp Ser Cys Arg Val Thr Lys Glu Asn Ser His Gln Val Lys Glu Glu  
195 200 205

Leu Ser Gln Lys Phe Asp Thr Leu Tyr Ala Ile Leu Asp Glu Lys Lys  
210 215 220

Ser Glu Leu Leu Gln Arg Ile Thr Gln Glu Gln Glu Glu Lys Leu Gly  
225 230 235 240

Phe Ile Glu Ala Leu Ile Leu Gln Tyr Arg Glu Gln Leu Glu Lys Ser  
245 250 255

Thr Lys Leu Val Glu Thr Ala Ile Gln Ser Leu Asp Glu Pro Gly Gly  
260 265 270

Ala Thr Phe Leu Ser Ser Ala Lys Gln Leu Ile Lys Ser Ile Val Glu  
 275 280 285

Ala Ser Lys Gly Cys Gln Leu Gly Lys Thr Glu Gln Gly Phe Glu Asn  
 290 295 300

Met Asp Tyr Phe Thr Leu Asp Leu Glu His Ile Ala Glu Ala Leu Arg  
 305 310 315 320

Ala Ile Asp Phe Gly Thr Gly Lys Gly Cys Asp Val Thr Cys Leu Thr  
 325 330 335

Phe Glu Arg Gln Arg Ser Ser  
 340

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